

# SEQUENCE LISTING

<110> James Karras  
Erich Koller

<120> ANTISENSE MODULATION OF TOLL-LIKE RECEPTOR 4 EXPRESSION

<130> ISPH-0618

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Met Glu Leu Asn  
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Ile Glu Asp Gly Ala Tyr Gln Ser Leu Ser His Leu Ser Thr Leu Ile	
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Ile Asp Leu Phe Asn Cys Leu Thr Asn Val Ser Ser Phe Ser Leu Val							
		265		270		275	
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Ser Val Thr Ile Glu Arg Val Lys Asp Phe Ser Tyr Asn Phe Gly Trp							
		280		285		290	
caa cat tta gaa tta gtt aac tgt aaa ttt gga cag ttt ccc aca ttg	1208						
Gln His Leu Glu Leu Val Asn Cys Lys Phe Gly Gln Phe Pro Thr Leu							
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Asn Ala Phe Ser Glu Val Asp Leu Pro Ser Leu Glu Phe Leu Asp Leu							
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Ser Arg Asn Gly Leu Ser Phe Lys Gly Cys Cys Ser Gln Ser Asp Phe							
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Gly Thr Thr Ser Leu Lys Tyr Leu Asp Leu Ser Phe Asn Gly Val Ile							
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acc atg agt tca aac ttc ttg ggc tta gaa caa cta gaa cat ctg gat	1448						
Thr Met Ser Ser Asn Phe Leu Gly Leu Glu Gln Leu Glu His Leu Asp							
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ttc cag cat tcc aat ttg aaa caa atg agt gag ttt tca gta ttc cta	1496						
Phe Gln His Ser Asn Leu Lys Gln Met Ser Glu Phe Ser Val Phe Leu							
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Ser Leu Arg Asn Leu Ile Tyr Leu Asp Ile Ser His Thr His Thr Arg							
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Val Ala Phe Asn Gly Ile Phe Asn Gly Leu Ser Ser Leu Glu Val Leu							
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aaa atg gct ggc aat tct ttc cag gaa aac ttc ctt cca gat atc ttc	1640						
Lys Met Ala Gly Asn Ser Phe Gln Glu Asn Phe Leu Pro Asp Ile Phe							
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aca gag ctg aga aac ttg acc ttc ctg gac ctc tct cag tgt caa ctg	1688						
Thr Glu Leu Arg Asn Leu Thr Phe Leu Asp Leu Ser Gln Cys Gln Leu							
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gag cag ttg tct cca aca gca ttt aac tca ctc tcc agt ctt cag gta	1736						
Glu Gln Leu Ser Pro Thr Ala Phe Asn Ser Leu Ser Ser Leu Gln Val							
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cta aat atg agc cac aac aac ttc ttt tca ttg gat acg ttt cct tat	1784						

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Leu	Asn	Met	Ser	His	Asn	Asn	Phe	Phe	Ser	Leu	Asp	Thr	Phe	Pro	Tyr	
485					490					495					500	
aag	tgt	ctg	aac	tcc	ctc	cag	gtt	ctt	gat	tac	agt	ctc	aat	cac	ata	1832
Lys	Cys	Leu	Asn	Ser	Leu	Gln	Val	Leu	Asp	Tyr	Ser	Leu	Asn	His	Ile	
			505						510					515		
atg	act	tcc	aaa	aaa	cag	gaa	cta	cag	cat	ttt	cca	agt	agt	cta	gct	1880
Met	Thr	Ser	Lys	Lys	Gln	Glu	Leu	Gln	His	Phe	Pro	Ser	Ser	Leu	Ala	
			520					525						530		
ttc	tta	aat	ctt	act	cag	aat	gac	ttt	gct	tgt	act	tgt	gaa	cac	cag	1928
Phe	Leu	Asn	Leu	Thr	Gln	Asn	Asp	Phe	Ala	Cys	Thr	Cys	Glu	His	Gln	
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Ser	Phe	Leu	Gln	Trp	Ile	Lys	Asp	Gln	Arg	Gln	Leu	Leu	Val	Glu	Val	
	550					555					560					
gaa	cga	atg	gaa	tgt	gca	aca	cct	tca	gat	aag	cag	ggc	atg	cct	gtg	2024
Glu	Arg	Met	Glu	Cys	Ala	Thr	Pro	Ser	Asp	Lys	Gln	Gly	Met	Pro	Val	
565					570					575					580	
ctg	agt	ttg	aat	atc	acc	tgt	cag	atg	aat	aag	acc	atc	att	ggg	gtg	2072
Leu	Ser	Leu	Asn	Ile	Thr	Cys	Gln	Met	Asn	Lys	Thr	Ile	Ile	Gly	Val	
				585					590					595		
tcg	gtc	ctc	agt	gtg	ctt	gta	gta	tct	gtt	gta	gca	gtt	ctg	gtc	tat	2120
Ser	Val	Leu	Ser	Val	Leu	Val	Val	Ser	Val	Val	Ala	Val	Leu	Val	Tyr	
			600					605					610			
aag	ttc	tat	ttt	cac	ctg	atg	ctt	ctt	gct	ggc	tgc	ata	aag	tat	ggg	2168
Lys	Phe	Tyr	Phe	His	Leu	Met	Leu	Leu	Ala	Gly	Cys	Ile	Lys	Tyr	Gly	
		615					620					625				
aga	ggg	gaa	aac	atc	tat	gat	gcc	ttt	gtt	atc	tac	tca	agc	cag	gat	2216
Arg	Gly	Glu	Asn	Ile	Tyr	Asp	Ala	Phe	Val	Ile	Tyr	Ser	Ser	Gln	Asp	
	630					635					640					
gag	gac	tgg	gta	agg	aat	gag	cta	gta	aag	aat	tta	gaa	gaa	ggg	gtg	2264
Glu	Asp	Trp	Val	Arg	Asn	Glu	Leu	Val	Lys	Asn	Leu	Glu	Glu	Gly	Val	
645					650					655					660	
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